List of relevant sequences

SEQ ID NO:1 (AJ002425.2 Nucleotide sequence for CAA05410.2 (LBDG5) protein, coding region start position =61, coding region finish position=1905)

```
1 ggggcttatt aacggtggta tctggtagag cccccataaa gtaccaccgt ggggaagaca
5
       61 atqqqacccq aaaacqaqaq ccccqtqqqq ggggagatgt gggggagagt gtgtgtgttc
      121 agcgctagac aaaagagacc gcgatgcagc actttaaata gtcttctcgt tgctagaaca
      181 tgctataaca tgctggcgct ggtggattgg atcgagcagg atttcaagag cacttggcga
      241 aaggeteacg ttteecteeg egggtteete gatttegage tggggagaeg aaaagaggtt
      301 gccggggctt ttctgggtgg ggataccaga cctgacccca aaaagccgcg gggggggtcc
10
      361 aaaaagaacg tggaggtgta tgatgatgac gtaggctctc aggctgcgga cagccccggg
      421 aaacqcatqq ccccqaaaqq gacatttaga gataaggaca aatttgaagg gctgtttaag
      481 ctcqqqqcqc tqqtqqcaaa aaaagccttg accccagcat tttcttgttc ccccaacagg
      541 gggtegeete tacaegeeca ttatggggat gaaateetet acaaggttga ateegggeee
15
      601 gtcaacattt gcgaggggg caaaagaggc gtggggatcc acccccaga taactacggc
      661 gataccettg atgaaaatet tggcetteee caaaaaattg tgattaaggt gaageeecaa
      721 accqaqqaaq ccaacacttq qttaaggcag gatctgaaaa atcataacag tgcaaaggag
      781 gccgggggct ccgacgagat taaaaccttt gtgacaggat gtaaaaaaga tgggcatagt
      841 gggcgtaaaa atatgaccac acatgacaga aattcaaaaa agtggcaacg ggtaaacttg
20
      901 tecttatgg ecteecteca getagattet aggggeggae gegeggggee eeggegegga
      961 gcgcggcgcc tgtgcctggt gtgtgaggac tatgccagct gttcaaacac ctgtgtctgg
     1021 tcctqtqaaq cctacaaqqt cttctttcgc cgaaqtcaaa gtttcacaga tccagcctgt
     1081 ttcacaaacq attqcaacat ctctaaqaat aggtctaagt cttgcccagc ttgcctcctc
     1141 cqttqcctqc accctaqcat taatqaqatc cqaaaaqaca agcgaqcaqc qctgaaaqtq
25
     1201 cqaqacaacq ttggtgaaga ggtggatatg accggtccta gctggacctg cctgaagcta
     1261 ctttttcaq atqqqqaaaa aqtqataccc agattgggcc atgaactccc agggatcaag
     1321 gggggccggc aggcaaaaca gcagtcccac cgaggaagcc ccatccccaa aaacaggaaa
     1381 qqttqqcccc ccqqacatqt cctqtcaaat gacqqcqqaq ctqqtqqcaq qqtatqqaaa
     1441 aaaaaatcct gtaaaccaat tcgccgagaa ggccccaagt ggtgggatcg gctgaatgaa
     1501 tctacacctt tgttttgggg gtctcgagcc aacaagagtt tagggaaggg aggcaccagg
30
     1561 gggaggattt tcatcaagca cccacacctc tttaagtttg cagcagatcc tcaggacaag
     1621 cactggctgg ctgagcagca tcatatgcgg gcaacaggag gaaagatggc gtaccttctc
     1681 attgaggaag acatcgggca gcatcatggc caggggttcc cagttatgct tctcaagatt
     1741 agccatatta ggcacatggt tgggggagtg gctcattgct tgtacgacat gaaagaaaag
     1801 aagtttgttc tgccatcctg gaaggttgag aagttgggga aatacgtgga gacactacgg
35
     1921 tqaqqccatc atggggatgc gqtctaqttg gctcttagca gcatcaagct gtacatgagc
      1981 tagtttgtag tgactcactg cagagccccc cagactggct tgtggttctg tttctaaagt
```

2041 tattggaata agaagcaatt aaacaagttt gtaatttaaa aaaaaaaaa aaaaa

SEQ ID NO:2 (Protein CAA05410.2; LBDG5)

1 mgpenespvg gemwgrvcvf sarqkrprcs tlnsllvart cynmlalvdw ieqdfkstwr
61 kahvslrgfl dfelgrrkev agaflggdtr pdpkkprggs kknvevyddd vgsqaadspg
121 krmapkgtfr dkdkfeglfk lgalvakkal tpafscspnr gsplhahygd eilykvesgp
181 vniceggkrg vgihppdnyg dtldenlglp qkivikvkpq teeantwlrq dlknhnsake
241 aggsdeiktf vtgckkdghs grknmtthdr nskkwqrvnl slmaslqlds rggragprrg
301 arrlclvced yascsntcvw sceaykvffr rsqsftdpac ftndcniskn rskscpacll
361 rclhpsinei rkdkraalkv rdnvgeevdm tgpswtclkl lfsdgekvip rlghelpgik
421 ggrqakqgh rgspipknrk gwppghvlsn dggaggrvwk kksckpirre gpkwwdrlne
481 stplfwgsra nkslgkggtr grifikhphl fkfaadpqdk hwlaeqhhmr atggkmayll
541 ieedigqhhg qgfpvmllki shirhmvggv ahclydmkek kfvlpswkve klgkyvetlr
601 tekehraaea spqt

15

20

25

30

35

10

5

SEQ ID NO:3 (Nucleotide coding sequence for LBDG10, CAA05409.2)

1 atggaccccg agaacgtgag agccgagggc ggggacatga gggagaaagt gagactgtcc 61 agcqctagac agagactgcg atgcagcact aagaatagtc ttctcgttgc tagaacatgc 121 gtgctgaggc tggtcgattg gatcgagcag gatctcaaga gcacttgcga aaaggctcac 181 gtttccctcc gcgggtacct cgatttccag ctgcaatgcg ggagacgaaa agaggttgcc 241 ggggctctgg gtggggatac cagacctgac cccaagaagc cgcggggggg ctccaaaaag 301 aacgtggagg tgtatggtga tgacgtaggc tctcaggctg cggacagccc gaggaaacag 361 ctqqccqcqa aaqqqacatt cagagataag gacaaaattg aagcgctgtt caagctcggg 421 gagctggtgg ctaaaaaagc cttgtcctca gcaattactt ggttccccaa cagcgtgtcg 481 cctctacacg cccattatgg ggatgaaatc ctctacaagg atgaatccgg gcttgtcaac 541 attagcgagg gtggcaaaag aggcgtggag atccacccc cagataactt cggcatcaca 601 accettgatg aagatettgg ettteeceaa ataattgtga ttaaegtgaa geeteaaace 661 gaggaagcca acacttggta tagacaggat ctgaaatatc ataacagtgc aaacgaggcc 721 qqqtactccq acgagaataa aacctttgtg agaggatgta gaaaagatgg gcatagtgag 781 cqtaacaata tgaccacagg tgacagaaat tcaaaaaagg cccagcccgt aaacttctcc 841 ctcatggcat ccctcgccct ggattctagg ggcaaagccg cgggggccccg gcgcggagcg 901 aggcgcctgt gcctggtgtg tgaggactat gccagctgtt caaacacctg tgtctggtcc 961 tgtgaageet acaaggtett etttegeega agteaaagtt teacagatee ageetgttte 1021 acaaacgatt gcaacatctc taagaataga tctaagtctt gcccagcctg cctcctccgt 1081 tgcctgcagc ctagcatcaa tgagatccga aaagacaagc gagcagcgct gaatgtgcga 1141 gacaacgttg gtgaagaggt ggatatgacc ggtcctagct ggacctgcct gaagctactc 1201 ttttcagatg gagaaaaagt gatacccaga ttggcccatg aacttccagg gatcaagcgt 1261 ggccggcagg cagaagagga gtcccaccga ggaagcccca ttcccaaaaa gaggaaaggt

tggcctcctg gacatgtcct gtcaaatgac cgcgcagctg ctggcacggt atggaaacca aatcctgtg aaccaattcg ccgagaaggc cccaagtggg acgctcggct gaatgaatct t441 accacctttg ttttggggtc tcgagccaac aaggccttag ggaagggagg caccagaggg 1501 aggatttaca tcaagcaccc acacctcttt aagtatgcag cagatcctca ggacaagcac 1561 tggctggctg agcagcttca tatgcgggca acaggaggaa agatggcgta ccttctcatt 1621 gaggaagaca tcgagcagca tcatggccag cggttcgcag ttttccttct caagattagc 1681 catattaggc acatggttga gggagtggct cattgcttgt acgacatgaa agttatccag 1741 tttgttctgc catcctggaa ggttgagaag ttgcggaaat acgtggagac actacggaca 1801 gaaaatgagc atcgtgctgc tgaagcaagt ccccagacct ga

10

5

SEQ ID NO:4 (Protein coding sequence for LBDG10, CAA05409.2)

1 mdpenvraeg gdmrekvrls sarqrlrcst knsllvartc vlrlvdwieg dlkstcekah
61 vslrgyldfq lqcgrrkeva galggdtrpd pkkprggskk nvevygddvg sqaadsprkq
15 121 laakgtfrdk dkiealfklg elvakkalss aitwfpnsvs plhahygdei lykdesglvn
181 iseggkrgve ihppdnfgit tldedlgfpq iivinvkpqt eeantwyrqd lkyhnsanea
241 gysdenktfv rgcrkdghse rnnmttgdrn skkaqpvnfs lmaslaldsr gkaagprrga
301 rrlclvcedy ascsntcvws ceaykvffrr sqsftdpacf tndcnisknr skscpacllr
361 clqpsineir kdkraalnvr dnvgeevdmt gpswtclkll fsdgekvipr lahelpgikr
20 421 grqaeeeshr gspipkkrkg wppghvlsnd raaagtvwkp kscepirreg pkwdarlnes
481 ttfvlgsran kalgkggtrg riyikhphlf kyaadpqdkh wlaeqlhmra tggkmaylli
541 eedieqhhgq rfavfllkis hirhmvegva hclydmkviq fvlpswkvek lrkyvetlrt
601 enehraaeas pqt

25 SEQ ID NO:5 (Nucleotide coding sequence for BAB62888.1 (LBDG11)

1 atggagcca ctggcgacgc cgagcagccg cggggaccca gcggggccga gaggggcggc
61 ttggagctgg gggatgcggg cgcagcgggg cagctggttc ttacgaaccc ttggaacata
121 atgataaagc accggcaggt gcagcggagg ggccgccgct cacagatgac aacaagtttc
181 acagatcctg ccatctccat ggatctcctc cgagctgtcc tgcagcccag catcaacgag
30 241 gagatccaga ctgtcttcaa caagtacatg aagttcttcc agaaggcagc actgaacgtg
301 cgagacaatg ttggggagga ggtggacgca gagcagctga tccaggaagc ctgtcggagc
361 tgcctggagc aggctaaact gctctttca gatggagaa aagtaatacc cagattgacc
421 catgagcttc caggaataaa gcgtggccgt caggcagaag aagaatgtgc ccatcgagga
481 agcccccttc ctaaaaagag gaaaggacgg cctcctggac acatcctgtc aagcgaccgg
35 541 gcagccgccg gcatggtatg gaaaccaaaa tcctgtgaac caattcgccg ggaaggccc
601 aagtgggacc cagctcgcct gaatgaatct accacctttg tgttgggatc tcgagccaac
661 aaagccctgg ggatggggg caccagagga agaatctaca tcaagcaccc acacctcttt
721 aagtatgcag ctgacccca ggataagcac tggctggctg agcagcatca catggggca

4

781 acaggggca agatggcta cctcctcatc gaggaggaca tccgggacct tgcggccagt 841 gatgattaca gaggatgcct ggatctgaag ctagaggaat tgaaatcctt tgtcctaccc 901 tcctggatgg tggagaagat gagaaagtat atggagacac tacggacaga gaatgagcat 961 cgtgctgttg aagcacctcc acagacctga

5

10

WO 2004/005344

SEO ID NO:6 (Protein coding sequence for BAB62888.1 (LBDG11)

1 mgatgdaeqp rgpsgaergg lelgdagaag qlvltnpwni mikhrqvqrr grrsqmttsf 61 tdpaismdll ravlqpsine eiqtvfnkym kffqkaalnv rdnvgeevda eqliqeacrs 121 cleqakllfs dgekviprlt helpgikrgr qaeeecahrg splpkkrkgr ppghilssdr 181 aaagmvwkpk scepirregp kwdparlnes ttfvlgsran kalgmggtrg riyikhphlf 241 kyaadpqdkh wlaeqhhmra tggkmaylli eedirdlaas ddyrgcldlk leelksfvlp 301 swmvekmrky metlrteneh raveappqt

SEQ ID NO:7 (Nucleotide coding sequence for AAH03486.1 (LBDG12)

15 1 atqqqqqcqa ctqqcqacac cgagcagccg cggggccccg gcggggcgga gcgaggtggc 61 ctqqaqctqq qcqacqcggg cgcggggc cagccggttc tcacgaaccc ttggaacata 121 atgataaaac atcggcaggt gcagcgaagg ggccgccgat ctcagatgac cacaagtttc 181 acagacccag ccatctctat ggatctcctc cgtgctgtcc tgcagcctag catcaatgag 241 gagatccagg gtgtcttcaa caagtacatg aagttcttcc agaaggcagc gctgaatgtg 301 cgagacaatg ttggggaaga agtggacgca gagcagttga ttcaggaggc ctgccgcagc 20 361 tgcctggagc aggcaaagct gctcttttct gatggagaga aagtgatacc cagattggcc 421 catgagette cagggateaa geggggeegg caageagaag aggagteeca eegaggaage 481 cccattccca aaaagaggaa aggtcggcct cctggacacg tcctgtcaaa tgaccgcgca 541 gctgctggca tggtatggaa acaaaaatcc tgtgaaccaa ttcgccgaga aggccccaag 601 tgggacccag ctcggctgaa tgaatctacc acctttgttt tggggtctcg agccaacaag 25 661 gccttgggga tgggaggcac cagagggaga atctacatca agcacccaca cctctttaag 721 tatqcaqcaq atcctcagga caagcactgg ctggctgagc agcatcacat gcgggcaaca 781 ggcggaaaga tggcgtacct tctcattgaa gaagacatcc gggacttggc tgccagcgat 841 gactacagag gatgettgga cetgaagttg gaggagetga aateetttgt tetgecatee 901 tggatggttg agaagatgcg gaaatacatg gagacactgc ggacagaaaa tgagcaccgc 30 961 gctgcggaag cgcctcccca gacctga

SEQ ID NO:8 (Protein coding sequence for AAH03486.1 (LBDG12)

1 mgatgdteqp rgpggaergg lelgdagaag qpvltnpwni mikhrqvqrr grrsqmttsf
35 61 tdpaismdll ravlqpsine eiggvfnkym kffqkaalnv rdnvgeevda eqliqeacrs
121 cleqakllfs dgekviprla helpgikrgr qaeeeshrgs pipkkrkgrp pghvlsndra
181 aagmvwkqks cepirregpk wdparlnest tfvlqsrank algmggtrgr iyikhphlfk

WO 2004/005344

5

241 yaadpqdkhw laeqhhmrat ggkmayllie edirdlaasd dyrgcldlkl eelksfvlps 301 wmvekmrkym etlrtenehr aaeappqt

SEQ ID NO:9 (Nucleotide coding sequence for AAK49953.1 (LBDG13)

1 atqqqqqqa ctqqcqacac cqgqqqnccc cqgccgggga cggaatcaag gcggcctgga 5 61 aacqtqqqca acqcqqqcqc qqcqqqccag ccggttctca cgaacccttg gaacataatg 121 atcaaacctc ggcaggtgca gcgaagaggc cgccgatctc agatgaccac aagtttcaca 181 gatccagcca tctctatgga tctcctccgt gctgtcctgc agcctagcat caatgaggag 241 atccagagtg tcttcaacaa gtacatgaag ttcttccaga aggcagcgct gaatgtgcga 301 gacaacgttg gtgaagaggt ggatgcagaa cagttgattc aggaggcctg ccgcagctgc 10 361 gtggagcagg caaagctact cttttcagat ggagaaaaag tgatacccag attggcccat 421 gaacttccag ggatcaagcg tggccggcag gcagaagagg agtcccaccg agaagcccca 481 ttcccaaaaa gaggaaaggt cggccttcct ggacatgtcc tgtcaaatga ccgcgcagct 541 gctggcatgg tatggaaacc aaaatcctgt gaaccaattc gccgagaagg ccccaagtgg 601 gaccagete ggetgaatga atetaceace tttgttttgg ggtetegage caacaaggee 15 661 ttagggatgg gaggcaccag agggaggatt tacatcaagc acccacacct ctttaagtat 721 gcagcagatc ctcaggacaa gcactggctg gctgagcagc atcatatgcg ggcaacagga 781 ggaaagatgg cgtaccttct cattgaggaa gacatccgag acttggctgc cagcgatgac 841 tacagaggat gcttggacct gaagttggag gagctgaagt cctttgtttt gccatcctgg 901 atggttgaga agatgcggaa atacatggag acactacgga cagaaaatga gcatcgtgct 20 961 gctgaagcaa ctccccagac ctga

SEQ ID NO:10 (Protein coding sequence for AAK49953.1 (LBDG13)

1 mgatgdtggp rpgtesrrpg nvgnagaagq pvltnpwnim ikprqvqrrg rrsqmttsft
61 dpaismdllr avlqpsinee iqsvfnkymk ffqkaalnvr dnvgeevdae qliqeacrsc
121 veqakllfsd gekviprlah elpgikrgrq aeeeshreap fpkrgkvglp ghvlsndraa
181 agmvwkpksc epirregpkw dparlnestt fvlgsranka lgmggtrgri yikhphlfky
241 aadpqdkhwl aeqhhmratg gkmaylliee dirdlaasdd yrgcldlkle elksfvlpsw
301 mvekmrkyme tlrtenehra aeatpgt

30

25